



REFERENCE LISTING

Chinketo, Richard A
Prayaga, Subhinder K

<1> Novel Humanized Brown Rat and Human Antic
Rhesusins Name

<11> 1544-117 PAF-X

<141> USNM 08/44,545

<141> 2000-01-31

<150> USNM 60/141,899

<151> 1999-07-27

<163> 13

<170> PatentIn Ver. 2.1

<210> 1

<211> 633

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

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Met Ala Pro Leu Ala Ala Val His Gly Ile Leu Gly His Ile His Gly

1

5

10

15

tta ggc acc tta gag ttc tta ggc acc tta gag ttc tta gag ttc 8*

Ile His Val His Val His Val His Val His Val His Val His Val His

acc acc acc acc acc acc acc acc acc acc acc acc acc acc acc 12*

Arg Ile Ile Leu Leu His His Arg Arg Arg Arg Arg Arg Arg Arg Arg

4

8

acc acc acc acc acc acc acc acc acc acc acc acc acc acc acc 16*

Arg His His Ile His Ala Ala Ile Ile Ala His Ile Ile Ile Ile Ile

Leu Gly Ala Ala Val Gly Ser His His Leu Leu His His Ala Gly Val
121 122 123 124 125 126 127 128 129 130 131 132 133 134 135 136 137 138 139 140

Arg His His Leu Leu Gly His Arg Arg Ser Ala Ala Val Arg Ser Ala
141 142 143 144 145 146 147 148 149 150 151 152 153 154 155 156 157 158 159 160

Arg Gly Gly His Gly Ala Ala His Leu Ala His Leu His Gly His His
161 162 163 164 165 166 167 168 169 170 171 172 173 174 175 176 177 178 179 180

Arg Arg Arg Gly Leu Tyr Tyr Arg Thr Gly His His Leu His His His
181 182 183 184 185 186 187 188 189 190 191 192 193 194 195 196 197 198 199 200

Pro Asp Gly Ser Val His Gly Thr Arg His Arg His Ser Leu His Gly
201 202 203 204 205 206 207 208 209 210 211 212 213 214 215 216 217 218 219 220

Ile Leu Glu Phe Ile Ser Val Ala Val Gly Leu Val Ser Ile Arg Gly
221 222 223 224 225 226 227 228 229 230 231 232 233 234 235 236 237 238 239 240

Val Asp Ser Gly Leu Tyr Leu Gly Met Asn Asp Lys Gly His Leu Tyr
241 242 243 244 245 246 247 248 249 250 251 252 253 254 255 256 257 258 259 260

Gly Ser Glu Lys Leu Thr Ser Glu Cys His Phe Arg His His Phe His
261 262 263 264 265 266 267 268 269 270 271 272 273 274 275 276 277 278 279 280

Glu Asn Trp Tyr Asn Thr Tyr Ser Ser Asn His Tyr Lys His Gly Asp
281 282 283 284 285 286 287 288 289 290 291 292 293 294 295 296 297 298 299 300

Thr Gly Arg Arg Tyr Phe Val Ala Leu Asn Lys Asp Gly Thr Phe Arg
301 302 303 304 305 306 307 308 309 310 311 312 313 314 315 316 317 318 319 320

Asp Gly Ala Arg Ser Lys Arg His Gln Lys Phe Thr His Phe Leu His
321 322 323 324 325 326 327 328 329 330 331 332 333 334 335 336 337 338 339 340

Arg Pro Val Asp Pro Glu Arg Val Pro His Leu Tyr Lys Arg Leu Leu
341 342 343 344 345 346 347 348 349 350 351 352 353 354 355 356 357 358 359 360

Val Tyr His
361 362 363

Val Tyr
364 365
366 367
368 369

<410> 4

ctggtaatt ctatattata gttatattat ctatattat

44

<210> 4

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<212> DNA

<213> Artificial Sequence

<210>

<213> Description of Artificial Sequence: primer

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ctgttctctg agagtgtaca tttataggtc ctta

45

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<212> DNA

<213> Homo sapiens

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tagaaatatt cccccattggt actatccagg gaaccaggaa agacttttgc gattttggt 120
ttctggaatt tctcagtata gttgtgggac tggtcagcat ttgcttgaat gacattttgc 180
tctacctcgg gatgaatgag aaggggggagc tgtatggatc agaaaaacta acccaagagt 240
gtgtattcag aqaacagttc gaagaaaaact ggtataatac gtaactgtca aacctatata 300
agcaagtgga caatggaagg cgtatctatg ttgtatttaa taattttgg attttttg 360
aagggattat gactaaaagg ctttttaatt tctatcattt ttatctaga ctatcttt 420
ctta 480

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<212> DNA

<213> Homo sapiens

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ctgttctctg agagtgtaca tttataggtc ctta
ctgttctctg agagtgtaca tttataggtc ctta
ctgttctctg agagtgtaca tttataggtc ctta

Figure 1. The effect of the number of trials on the number of correct responses. The number of correct responses was significantly higher for the 10-trial condition than for the 5-trial condition. Error bars represent the standard error of the mean.

[illegible]

1. *Phragmites* (1990)

• 213> Homo sapiens

ttgagtcata	gagttatcct	ttgtttatca	tttcaagata	gagatcatt	tttttttttt	68
taatactgaa	gagtcctcct	gcctacactga	ctttattttaa	tataatt		10

011> 308

213> How sapiens:

Val Pro Phe Gly Asn Val Thr Val Leu Pro Val Asp Thr Pro Val Ser

Leu Ser Asp His Leu Gly Gln Ser Glu Ala Gly Gly Leu Pro Arg Gly

Gly Leu Tyr Leu Gly Met Asn Glu Lys Tyr Glu Leu Tyr Tyr Ser Thr
 117 118 119

Lys Leu Thr Ala Glu Lys Val Phe Asn Glu Glu Phe Glu Glu Asn Lys
 120 121 122

Tyr Asn Thr Tyr Ser Ser Asn Leu Tyr Lys His Val Asp Thr Tyr Asn
 123 124 125

Asn Tyr Tyr Val Ala Leu Asn Lys Asp Lys Thr His Asn Ser Gly Thr
 126 127 128

Asn Thr Lys Asn His Glu Lys Phe Thr His His Leu His Asn His Val
 129 130 131

Asp Pro Asp Lys Val Pro Glu Leu Tyr Lys Asp His Leu Ser Glu Ser
 132 133 134

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<211> 208

<212> PRT

<213> Mus sp.

<400> 10

Met Ala Pro Leu Gly Glu Val Gly Ser Tyr Phe Gly Val Glu Asp Ala
 1 2 3 4 5 6 7 8 9 10

Val Pro Phe Gly Asn Val Pro Val Leu Pro Val Asp Ser His Val Leu
 11 12 13 14 15 16 17 18 19 20

Leu Asn Asp His Leu Gly His Ser Glu Asn Thr Tyr Leu Thr Tyr
 21 22 23 24 25 26 27 28 29 30

Asn Val Ser Thr Asp Leu Asn His Leu Lys Glu Thr Ser Asn Thr Thr
 31 32 33 34 35 36 37 38 39 40

Glu Leu Tyr Tyr Asn Thr Lys Phe His Leu His His His His Asn Tyr
 41 42 43 44 45 46 47 48 49 50

His Leu Ser Lys Thr Asn Thr Asn His Thr Thr Thr Thr Thr Thr

